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OM protein - protein search, using sw model

Run on: December 5, 2003, 18:14:44 ; Search time 28.64 Seconds
(without alignments)
264.443 Million cell updates/sec

Title: US-10-084-298-2

Perfect score: 903

Sequence: 1 MAALQKSVSSPLMGLTATSC.....EIKAIIGELDILFMSLRNACI 179

Scoring table: BILOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 328717 seqs, 42310858 residues

Total number of hits satisfying chosen parameters: 328717

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database : Issued_Patents_MA:
1: /cgn2_6_ptodata/2/iaa/5A_COMBO.pep:
2: /cgn2_6_ptodata/2/iaa/5B_COMBO.pep:
3: /cgn2_6_ptodata/2/iaa/6A_COMBO.pep:
4: /cgn2_6_ptodata/2/iaa/6B_COMBO.pep:
5: /cgn2_6_ptodata/2/iaa/PCTMUS_COMBO.pep:
6: /cgn2_6_ptodata/2/iaa/backTles1.pep:
*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	903	100.0	179	4	US-09-870-574-2
2	881	97.6	179	4	US-09-419-568F-28
3	881	97.6	179	4	US-09-354-243B-18
4	724	80.2	179	3	US-09-178-973B-15
5	724	80.2	179	4	US-09-419-568F-27
6	724	80.2	179	4	US-09-354-243B-27
7	720	79.7	179	3	US-09-178-973B-16
8	109	12.1	170	1	US-08-270-805C-2
9	109	12.1	170	2	US-08-410-54B-2
10	109	12.1	170	2	US-08-474-851-2
11	109	12.1	170	2	US-08-481-560-2
12	109	12.1	170	2	US-08-934-959-4
13	109	12.1	170	3	US-08-170-113-2
14	109	12.1	170	3	US-08-765-094C-26
15	109	12.1	170	3	US-09-082-797-26
16	109	12.1	170	3	US-08-643-810A-2
17	109	12.1	170	4	US-09-552-613-2
18	104.8	11.6	154	4	US-09-452-624A-3
19	102.5	11.4	178	1	US-08-270-805C-1
20	102.5	11.4	178	2	US-08-410-654B-1
21	102.5	11.4	178	2	US-08-474-51-1
22	103.5	11.4	178	2	US-08-481-560-1
23	102.5	11.4	178	2	US-08-934-959-6
24	102.5	11.4	178	3	US-08-170-113-1
25	102.5	11.4	178	3	US-08-765-094C-25
26	102.5	11.4	178	3	US-09-082-797-25
27	102.5	11.4	178	3	US-08-643-810A-1

ALIGNMENTS

RESULT 1
US-09-870-574-2

Sequence 2, Application US/09870574

Patent No. 6557799

GENERAL INFORMATION

APPLICANT: Gurney, Austin L.

APPLICANT: Aggarwal, Sudeepa.

APPLICANT: Xie Ming-Hong

APPLICANT: Maruoka, Ellen M.

APPLICANT: Foster, Jessica S.

APPLICANT: Goddard, Audrey

APPLICANT: Wood, William T.

TITLE OF INVENTION: INTERLEUKIN-22 POLYPEPTIDES, NUCLEIC ACIDS ENCODING TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDERS

FILE REFERENCE: P2806-1 (US)

CURRENT APPLICATION NUMBER: US/09-006-1 (US)

CURRENT FILING DATE: 2001-05-30

PRIOR APPLICATION NUMBER: US 60/169,495

PRIOR FILING DATE: 1999-12-07

PRIOR APPLICATION NUMBER: PCT/US00/14042

PRIOR FILING DATE: 2000-05-22

PRIOR APPLICATION NUMBER: PCT/US00/23328

PRIOR FILING DATE: 2000-08-24

NUMBER OF SEQ ID NOS: 7

SEQ ID NO 2

LENGTH: 179

TYPE: PRT

ORGANISM: Homo Sapien

US-09-870-574-2

Query Match 100.0%; Score 903; DB 4; Length 179;

Best Local Similarity 100.0%; Pred. No. 1.9e-100;

Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAALQKSVSSPLMGLTATSC.....EIKAIIGELDILFMSLRNACI 179

Db 1 MAALQKSVSSPLMGLTATSC.....EIKAIIGELDILFMSLRNACI 179

Qy 61 KFASLADNTDVRLIGEKLFQGVMSMERCYLKQVLFNFTLVEVLFFQSDRFQPMQEVVP 120

Db 61 KFASLADNTDVRLIGEKLFQGVMSMERCYLKQVLFNFTLVEVLFFQSDRFQPMQEVVP 120

Qy 121 FLARLSNRSLSTCHIEGDDLHTIQRNVYKLGEGSEIKAIIGELDILFMSLRNACI 179

Db 121 FLARLSNRSLSTCHIEGDDLHTIQRNVYKLGEGSEIKAIIGELDILFMSLRNACI 179

RESULT 2
US-09-419-568F-28

Sequence 28, Application US/09419568F

Patent No. 6331613
 GENERAL INFORMATION:
 APPLICANT: Dumoutier, Laure
 APPLICANT: Louhidi, Jamila
 APPLICANT: Renaud, Jean-Christophe
 TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors (TIFs) the Proteins Encoded, and Uses Thereof
 FILE REFERENCE: LUD 5543.2
 CURRENT APPLICATION NUMBER: US/09/419,568F
 CURRENT FILING DATE: 1999-10-18
 PRIOR APPLICATION NUMBER: US09/354,243
 PRIOR FILING DATE: 1999-07-16
 PRIOR APPLICATION NUMBER: US09/178,973
 PRIOR FILING DATE: 1998-10-26
 NUMBER OF SEQ ID NOS: 29
 SEQ ID NO 28
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo sapiens
 FEATURE:
 US-09-419-568F-28

Query Match 97.6%; Score 881; DB 4; Length 179;
 Best Local Similarity 97.8%; Pred. No. 8.3e-98;
 Matches 175; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 MAALQKVSSFLMGTIATSCILLALIVQGGAAPISSHCRDKSNFOQPYITNRTFMLA 60
 Db 1 MAALQKVSSFLMGTIATSCILLALIVQGGAAPISSHCRDKSNFOQPYITNRTFMLA 60

Qy 61 KEASLADNNTDVRLIGKLFHGVSMSRVCYLMKQVLFNFTLEVLFFQSDRFQPYMQEVVP 120
 Db 61 KEASLADNNTDVRLIGKLFHGVSMSRVCYLMKQVLFNFTLEVLFFQSDRFQPYMQEVVP 120

Qy 121 FLARLSNRSLSTCHIEGDDLHQIQRNVQKLKDVTYKLGESGEIKAIGBDLIFMSLRNACI 179
 Db 121 FLARLSNRSLSTCHIEGDDLHQIQRNVQKLKDVTYKLGESGEIKAIGBDLIFMSLRNACI 179

RESULT 3
 US-09-354-243B-28
 Sequence 28, Application US/09354243B
 Parent No. 6359117
 GENERAL INFORMATION:
 APPLICANT: Dumoutier, Laure
 APPLICANT: Louhidi, Jamila
 APPLICANT: Renaud, Jean-Christophe
 TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors (TIFs) the Proteins Encoded, and Uses Thereof
 FILE REFERENCE: LUD 5543.1
 CURRENT APPLICATION NUMBER: US/09/354,243B
 CURRENT FILING DATE: 1999-07-16
 PRIOR APPLICATION NUMBER: US09/178,973
 PRIOR FILING DATE: 1998-10-26
 NUMBER OF SEQ ID NOS: 29
 SEQ ID NO 28
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo sapiens
 FEATURE:
 US-09-354-243B-28

Query Match 97.6%; Score 881; DB 4; Length 179;
 Best Local Similarity 97.8%; Pred. No. 8.3e-98;
 Matches 175; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 MAALQKVSSFLMGTIATSCILLALIVQGGAAPISSHCRDKSNFOQPYITNRTFMLA 60
 Db 1 MAALQKVSSFLMGTIATSCILLALIVQGGAAPISSHCRDKSNFOQPYITNRTFMLA 60

Qy 61 KEASLADNNTDVRLIGKLFHGVSMSRVCYLMKQVLFNFTLEVLFFQSDRFQPYMQEVVP 120
 Db 61 KEASLADNNTDVRLIGKLFHGVSMSRVCYLMKQVLFNFTLEVLFFQSDRFQPYMQEVVP 120

RESULT 4
 US-09-178-973B-15
 Sequence 15, Application US/09178973B
 Patient No. 6274710
 GENERAL INFORMATION:
 APPLICANT: Dumoutier, Laure
 APPLICANT: Louhidi, Jamila
 APPLICANT: Renaud, Jean-Christophe
 TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors (TIFs) the Proteins Encoded, and Uses Thereof
 FILE REFERENCE: LUD 5543
 CURRENT FILING DATE: 1998-10-26
 NUMBER OF SEQ ID NOS: 17
 SEQ ID NO 15
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Mus musculus
 US-09-178-973B-15

Query Match 80.2%; Score 724; DB 3; Length 179;
 Best Local Similarity 78.2%; Pred. No. 5.9e-79;
 Matches 140; Conservative 19; Mismatches 20; Indels 0; Gaps 0;

Qy 1 MAALQKVSSFLMGTIATSCILLALIVQGAAAPISSHCRDKSNFOQPYITNRTFMLA 60
 Db 1 MAVLQSMSSFLMGTIATSCILLALIVQGAAAPISSHCRDKSNFOQPYITNRTFMLA 60

Qy 61 KEASLADNNTDVRLIGKLFHGVSMSRVCYLMKQVLFNFTLEVLFFQSDRFQPYMQEVVP 120
 Db 61 KEASLADNNTDVRLIGKLFHGVSMSRVCYLMKQVLFNFTLEVLFFQSDRFQPYMQEVVP 120

Qy 121 FLARLSNRSLSTCHIEGDDLHQIQRNVQKLKDVTYKLGESGEIKAIGBDLIFMSLRNACI 179
 Db 121 FLTRKLSNQLSSCHISGDDQNIQRNVRLKETVKGESGEIKAIGBDLIFMSLRNACI 179

RESULT 5
 US-09-419-568F-27
 Sequence 27, Application US/09419568F
 Patient No. 6331613
 GENERAL INFORMATION:
 APPLICANT: Dumoutier, Laure
 APPLICANT: Louhidi, Jamila
 APPLICANT: Renaud, Jean-Christophe
 TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducible Factors (TIFs) the Proteins Encoded, and Uses Thereof
 FILE REFERENCE: LUD 5543.2
 CURRENT APPLICATION NUMBER: US/09/419,568F
 CURRENT FILING DATE: 1999-10-18
 PRIOR APPLICATION NUMBER: US09/354,243
 PRIOR FILING DATE: 1999-07-16
 PRIOR APPLICATION NUMBER: US09/178,973
 PRIOR FILING DATE: 1998-10-26
 NUMBER OF SEQ ID NOS: 29
 SEQ ID NO 27
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Mus musculus
 US-09-419-568F-27

Query Match 80.2%; Score 724; DB 4; Length 179;
 Best Local Similarity 78.2%; Pred. No. 5.9e-79;
 Matches 140; Conservative 19; Mismatches 20; Indels 0; Gaps 0;

Qy 1 MAALQKVSSFLMGTIATSCILLALIVQGGAAPISSHCRDKSNFOQPYITNRTFMLA 60
 Db 1 MAALQKVSSFLMGTIATSCILLALIVQGGAAPISSHCRDKSNFOQPYITNRTFMLA 60

Qy 61 KEASLADNNTDVRLIGKLFHGVSMSRVCYLMKQVLFNFTLEVLFFQSDRFQPYMQEVVP 120
 Db 61 KEASLADNNTDVRLIGKLFHGVSMSRVCYLMKQVLFNFTLEVLFFQSDRFQPYMQEVVP 120

RESULT 6
 Qy 1 MAALQVSSESLMGLATSCILLALIVQGAAAPISLSCBLDKSNFQPYITNRTFMLA 60
 Db 1 MAVLQSMSFSLMGLAASCLLIALWAQANALPVNTRCKLEYSNFQPYITNRTFMLA 60
 Qy 61 KEASLADNNTDVRLIGEKLPHGVMSERCLMKGVLNFTLLEYLPPQSDRFQPYMVEVP 120
 Db 61 KEASLADNNTDVRLIGEKLPHGVMSERCLMKGVLNFTLLEYLPPQSDRFQPYMVEVP 120
 Qy 121 FLARLSNRLSTCHIEGDDLHQRLKDTVKLGESGEIKAIGELDLMFLSRLNACI 179
 Db 121 FLTKLSNQLSSCHISGDDNQIQNVRKRLKETVKLGESGEIKAIGELDLMFLSRLNACI 179

RESULT 6
 US-09-354-243B-27
 ; Sequence 27, Application US/09354243B
 ; Patent No. 6359117
 ; GENERAL INFORMATION:
 ; APPLICANT: Dumoutier, Laure
 ; APPLICANT: Louhed, Jamila
 ; APPLICANT: Renaud, Jean-Christophe
 ; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducers
 ; TITLE OF INVENTION: (TIFs)
 ; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
 ; FILE REFERENCE: LUD 5543.1
 ; CURRENT APPLICATION NUMBER: US/09/354,243B
 ; CURRENT FILING DATE: 1999-07-16
 ; PRIOR APPLICATION NUMBER: US09/178,973
 ; PRIOR FILING DATE: 1998-10-26
 ; NUMBER OF SEQ ID NOS: 29
 ; SEQ ID NO: 27
 ; LENGTH: 179
 ; TYPE: PRT
 ; ORGANISM: Mus musculus
 ; FEATURE:
 US-09-354-243B-27

Query Match 80 2%; Score 724; DB 4; Length 179;
 Best Local Similarity 78.2%; Pred. No. 5.9e-79; Mismatches 20; Indels 0; Gaps 0;
 Matches 140; Conservative 19; Mismatches 20; Indels 0; Gaps 0;

Qy 1 MAALQVSSESLMGLATSCILLALIVQGAAAPISLSCBLDKSNFQPYITNRTFMLA 60
 Db 1 MAVLQSMSFSLMGLAASCLLIALWAQANALPVNTRCKLEYSNFQPYITNRTFMLA 60
 Qy 61 KEASLADNNTDVRLIGEKLPHGVMSERCLMKGVLNFTLLEYLPPQSDRFQPYMVEVP 120
 Db 61 KEASLADNNTDVRLIGEKLPHGVMSERCLMKGVLNFTLLEYLPPQSDRFQPYMVEVP 120
 Qy 121 FLARLSNRLSTCHIEGDDLHQRLKDTVKLGESGEIKAIGELDLMFLSRLNACI 179
 Db 121 FLTKLSNQLSSCHISGDDNQIQNVRKRLKETVKLGESGEIKAIGELDLMFLSRLNACI 179

RESULT 7
 US-09-178-973B-16
 ; Sequence 16, Application US/09178973B
 ; Patent No. 6274710
 ; GENERAL INFORMATION:
 ; APPLICANT: Dumoutier, Laure
 ; APPLICANT: Louhed, Jamila
 ; APPLICANT: Renaud, Jean-Christophe
 ; TITLE OF INVENTION: Isolated Nucleic Acid Molecules which Encode T Cell Inducers
 ; TITLE OF INVENTION: (TIFs)
 ; TITLE OF INVENTION: The Proteins Encoded, and Uses Thereof
 ; FILE REFERENCE: LUD 5543
 ; CURRENT APPLICATION NUMBER: US/09/178,973B
 ; CURRENT FILING DATE: 1998-10-26
 ; NUMBER OF SEQ ID NOS: 17
 ; SEQ ID NO: 16
 ; LENGTH: 179
 ; TYPE: PPT

ORGANISM: Mus musculus
US-09-178-973B-16

Query Match 7
Best Local Similarity 7
Matches 139; Conservati

Qy	1	MA10LQSVSSFM
Db	1	MAV1QKMSFSLM
Qy	61	KEASLADINTDVE
Db	61	KEASLADINTDVE
Qy	121	FLARLNSRBLSTCH
Db	121	FLTRLNSQLSSSCH

RESULT 8
US-08-270-805C-2

1 Sequence 2, Application U
; Patent No. 5716451

GENERAL INFORMATION:

1 APPLICANT: Di-Hwei Hs
; APPLICANT: Kevin K. M
; APPLICANT: Hergen Spie
; TITLE OF INVENTION: o
; TITLE OF INVENTION: o
; NUMBER OF SEQUENCES:
; CORRESPONDENCE ADDRESS:
; ADDRESSEES: Schering
; STREET: 2000 Gallo
; CITY: Kenilworth
; STATE: New Jersey
; COUNTRY: USA
; ZIP: 07033

COMPUTER READABLE FORM

1 MEDIUM TYPE: Floppy
; COMPUTER: Macintosh
; OPERATING SYSTEM: 7

CURRENT APPLICATION DA

1 APPLICATION NUMBER:
; FILING DATE: 05-jul
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE: 23-dec
; APPLICATION NUMBER:
; FILING DATE: 04-feb
; APPLICATION NUMBER:
; FILING DATE: 16-jan
; ATTORNEY/AGENT INFORMATION:
; NAME: Pouille, Cynth
; REGISTRATION NUMBER:
; SEQUENCE CHARACTERISTICS:
; REFERENCE/DOCKET NUM
; TELECOMMUNICATION INFO:
; TELEPHONE: 908-298-
; TELEFAX: 908-298-53

1 INFORMATION FOR SEQ ID N
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1170 amino acid
; TYPE: amino acid
; STRANDEDNESS: singl
; TOPOLOGY: linear
; MOLECULE TYPE: peptid

1 Query Match 1
Best Local Similarity 2
Matches 43; Conservati

US-08-270-805C-2

QY 20 CILLLALIVQGGAAPISSHCRLDKSNFQQ-----PYITNTRTFMLEAKASLADNNTDV 72
 Db 11 CLVLLYLAPECGG----TDQC----DNPQMLRDLRDAFSRKTFQTKD----EVDN 56
 QY 73 RLIGEKLPHGVSMSERCYLMQVNLNFTEELVLFQPSDRFQPMQEVVPPFLAR----LSNR 128
 Db 57 LLIKESLILDFKGYLGCOALSEMIFQYLFEEVM-PQAEQDPEAKDHNTSLGENLKTLRLR 115

QY 129 LSITCHIEDDHLIQRNQVQLKDTVKLGESEGEKTAISEBLLD 169
 Db 116 LRRCHRFPCENKSKAVEQKNAFNKLOEKIYKAMSEFDI 156

RESULT 9
 US-08-410-654B-2
 Sequence 2, Application US/08410654B
 Patent No. 5833732

GENERAL INFORMATION:
 APPLICANT: Rene de Waal Malefyt
 APPLICANT: Di-Hwei Hsu
 APPLICANT: Anne O'Garra
 APPLICANT: Herzen Spits
 TITLE OF INVENTION: Use of An Interleukin-10 Antagonist to Treat
 TITLE OF INVENTION: Use of An Interleukin-10 Antagonist to Treat
 NUMBER OF SEQUENCES: 61

CORRESPONDENCE ADDRESS:
 ADDRESSEE: Schering-Plough Corporation
 STREET: 2000 Galloping Hill Road
 CITY: Kenilworth
 STATE: New Jersey
 COUNTRY: USA
 ZIP: 07033

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: Macintosh
 OPERATING SYSTEM: 7.5.3
 SOFTWARE: Microsoft Word 6.0

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/410,654B
 FILING DATE: 07-MAR-1995
 CLASSIFICATION: 424
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 08/229,854
 FILING DATE: 19-APR-1994
 APPLICATION NUMBER: US 08/410,654B
 FILING DATE: 24-MAR-1995
 CLASSIFICATION: 424
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 08/229,854
 FILING DATE: 19-APR-1994
 APPLICATION NUMBER: US 07/926,853
 FILING DATE: 06-AUG-1991
 ATTORNEY/AGENT INFORMATION:
 NAME: Poulik, Cynthia L.
 REGISTRATION NUMBER: DX0221KQ1GD
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 908-298-5388
 TELEFAX: 908-298-2987
 INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 170 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: Linear
 MOLECULE TYPE: peptide
 US-08-410-654B-2

Query Match 12.1%; Score 109; DB 2; Length 170;
 Best Local Similarity 26.7%; Pred. No. 3.7e-05; Mismatches 68; Indels 26; Gaps 6;
 Matches 43; Conservative 24; N mismatches 68; Indels 26; Gaps 6;

QY 20 CILLLALIVQGGAAPISSHCRLDKSNFQQ-----PYITNTRTFMLEAKASLADNNTDV 72
 Db 11 CLVLLYLAPECGG----TDQC----DNPQMLRDLRDAFSRKTFQTKD----EVDN 56

QY 73 RLIGEKLPHGVSMSERCYLMQVNLNFTEELVLFQPSDRFQPMQEVVPPFLAR----LSNR 128
 Db 57 LLIKESLILDFKGYLGCOALSEMIFQYLFEEVM-PQAEQDPEAKDHNTSLGENLKTLRLR 115

QY 129 I-STCHIEGDDLHIQRNYQKLUKTVKLGSEGIKAIGELDL 169
 Db 116 LRRCHRFPCENKSRAVEQKNAFKNLQEGIYAMSEFDI 156

RESULT 11
 US-08-481-560-2
 Sequence 2, Application US/08481560
 ; Patent No. 5837293
 GENERAL INFORMATION:
 ; APPLICANT: Rene de Waal Malefyt
 ; APPLICANT: Di-Hwei Hsu
 ; APPLICANT: Anne O'Garran
 ; APPLICANT: Hergen Spits
 TITLE OF INVENTION: Use of Interleukin-10 to Modulate
 TITLE OF INVENTION: Inflammation or T-Cell Mediated
 NUMBER OF SEQUENCES: 61
 CORRESPONDENCE ADDRESS:
 STREET: 2000 Galloping Hill Road
 CITY: Kenilworth
 STATE: New Jersey
 COUNTRY: USA
 ZIP: 07033

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: Macintosh
 OPERATING SYSTEM: 7.5.3
 SOFTWARE: Microsoft Word 6.0
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/481,560
 FILING DATE: 07-JUN-1995
 CLASSIFICATION: 424
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/410,654
 FILING DATE: 24-MAR-1995
 APPLICATION NUMBER: US 08/229,854
 FILING DATE: 19-APR-1994
 APPLICATION NUMBER: US 07/926,853
 FILING DATE: 06-AUG-1992
 APPLICATION NUMBER: US 07/742,129
 FILING DATE: 06-AUG-1991
 ATTORNEY/AGENT INFORMATION:
 NAME: Poulik, Cynthia L.
 REFERENCE NUMBER: 32,364
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 908-298-5388
 TELEFAX: 908-298-2987
 INFORMATION FOR SEQ ID NO: 2:
 INFORMATION CHARACTERISTICS:
 LENGTH: 170 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-08-481-560-2

Query Match 12.1%; Score 109; DB 2; Length 170;
 Best Local Similarity 26.7%; Pred. No. 3.7e-05;
 Matches 43; Conservative 24; Mismatches 68; Indels 6;
 Gaps 6;
 SEQUENCE CHARACTERISTICS:
 LENGTH: 170 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide

QY 20 CILLIALLVQGGAAAPLSSHCRLDKSNFQ-----PYITITFMAKEASLADNTDV 72
 Db 11 CIVLVLYAAPCGG----TDQC----DNFQMLRDLRDAFSRVTFFQTKD----EVDN 56

RESULT 12
 US-08-934-959-4
 Sequence 4, Application US/08934959
 ; Patent No. 598967
 GENERAL INFORMATION:
 ; APPLICANT: Knappe, Andrea
 ; APPLICANT: Fickenscher, Helmut
 ; APPLICANT: Fleckenstein, Bernhard
 ; TITLE OF INVENTION: MAMMALIAN CYTOKINE; RELATED REAGENTS
 ; NUMBER OF SEQUENCES: 7
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: DNAX Research Institute
 ; STREET: 901 California Avenue
 ; CITY: Palo Alto
 ; STATE: California
 ; COUNTRY: USA
 ; ZIP: 94304-1104
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/934,959
 FILING DATE: 22-SEP-1997;
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 60/027,368
 FILING DATE: 23-SEP-1996
 ATTORNEY/AGENT INFORMATION:
 NAME: Ching, Edwin P.
 REGISTRATION NUMBER: 34,090
 REFERENCE/DOCKET NUMBER: DXD644K
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 650-496-1200
 TELEFAX: 650-485-9196
 INFORMATION FOR SEQ ID NO: 4:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 170 amino acids
 TYPE: amino acid
 STRANDEDNESS: not relevant
 TOPOLOGY: not relevant
 MOLECULE TYPE: peptide
 US-08-934-959-4

Query Match 12.1%; Score 109; DB 2; Length 170;
 Best Local Similarity 26.7%; Pred. No. 3.7e-05;
 Matches 43; Conservative 24; Mismatches 68; Indels 26; Gaps 6;
 QY 20 CILLIALLVQGGAAAPLSSHCRLDKSNFQ-----PYITITFMAKEASLADNTDV 72
 Db 11 CIVLVLYAAPCGG----TDQC----DNFQMLRDLRDAFSRVTFFQTKD----EVDN 56

RESULT 13
 US-08-170-113-2
 Sequence 13, Application US/08170113
 ; Patent No. 6106823
 GENERAL INFORMATION:
 ; APPLICANT: Vieira, Paulo J.
 ; APPLICANT: Moore, Kevin W.
 ; APPLICANT: de Waal Malefyt, Rene

APPLICANT: de Vries, Jan E.
 APPLICANT: Fluckinger, Anne-Catherine
 APPLICANT: Banchereau, Jacques
 TITLE OF INVENTION: TREATMENT OF NEOPLASTIC DISEASE WITH
 TITLE OF INVENTION: INTERLEUKIN-10
 NUMBER OF SEQUENCES: 17
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: DNAX Research Institute
 STREET: 901 California Avenue
 CITY: Palo Alto
 STATE: California
 COUNTRY: USA
 ZIP: 94304-1104

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent In Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/170,113
 FILING DATE: 17-DEC-1993
 CLASSIFICATION: 54
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 07/933,419
 FILING DATE: 21-AUG-1992
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 08/091,333
 FILING DATE: 12-JUL-1993
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 08/020,018
 FILING DATE: 17-FEB-1993
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 07/830,496
 FILING DATE: 04-FEB-1992
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 07/641,347
 FILING DATE: 16-JAN-1991
 ATTORNEY/AGENT INFORMATION:
 NAME: Ching, Edwin P.
 REGISTRATION NUMBER: 34,090
 REFERENCE/DOCKET NUMBER: DX0116Q1KX
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 415-852-9196
 TELEFAX: 415-491-1200
 INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 170 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US -08-170-113-2

Query 20 CILLLALVQGAAAPISSHCRLLDKVINFLEEVLYFQSDRFPYMOEVVPPFLAR----LSNR 128
 Best Local Similarity 12.1%; Score 109; DB 3; Length 170;
 Matches 43; Conservative 24; Mismatches 68; Indels 26; Gaps 6;

Query 73 RLIGEKLFPHGYSMRSERYLCKMVKVINFLEEVLYFQSDRFPYMOEVVPPFLAR----LSNR 128
 Best Local Similarity 26.7%; Freq. No. 3.7e-05; Length 170;
 Matches 57; Conservative 24; Mismatches 68; Indels 26; Gaps 6;

Query 129 LSTCHIEGDDLHIIQRNVQLKDTVKLGESESSEKIAIGELD 169
 Best Local Similarity 12.1%; Score 109; DB 3; Length 170;
 Matches 57; Conservative 24; Mismatches 68; Indels 26; Gaps 6;

Query 20 CILLLALVQGAAAPISSHCRLLDKVINFLEEVLYFQSDRFPYMOEVVPPFLAR----LSNR 128
 Best Local Similarity 12.1%; Score 109; DB 3; Length 170;
 Matches 43; Conservative 24; Mismatches 68; Indels 26; Gaps 6;

Query 73 RLIGEKLFPHGYSMRSERYLCKMVKVINFLEEVLYFQSDRFPYMOEVVPPFLAR----LSNR 128
 Best Local Similarity 26.7%; Freq. No. 3.7e-05; Length 170;
 Matches 57; Conservative 24; Mismatches 68; Indels 26; Gaps 6;

Query 129 LSTCHIEGDDLHIIQRNVQLKDTVKLGESESSEKIAIGELD 169
 Best Local Similarity 12.1%; Score 109; DB 3; Length 170;
 Matches 57; Conservative 24; Mismatches 68; Indels 26; Gaps 6;

RESULT 15
 US -09-082-797-26
 Sequence 26, Application US/09082797

Patent No. 6168791
 GENERAL INFORMATION:
 APPLICANT: GRONHOU LARSEN, Christian
 APPLICANT: GESSER, Borbala
 TITLE OF INVENTION: IMMUNOMODULATORS
 NUMBER OF SEQUENCES: 27

CORRESPONDENCE ADDRESS:
 ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
 STREET: 419 Seventh Street N.W., Suite 300
 CITY: Washington

RESULT 14
 US -08-765-094C-26
 Sequence 26, Application US/08765094C

STATE: D.C.
 COUNTRY: U.S.A.
 ZIP: 20004
 COMPUTER READABLE FORM:
 COMPUTER: IBM PC compatible
 MEDIUM TYPE: Floppy disk
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/082,797
 FILING DATE:
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: 08/765,094
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: DK 0800/94
 FILING DATE: 05-JUL-1994
 ATTORNEY/AGENT INFORMATION:
 NAME: COOPER, Iver P.
 REGISTRATION NUMBER: 28,005
 REFERENCE/DOCKET NUMBER: GRONHOU-LARSEN=1
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (202) 628-5197
 TELEFAX: (202) 737-3528
 INFORMATION FOR SEQ ID NO: 26:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 170 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-09-082-797-26

Qy 20 CILILALYQGAAAPISSHORLDSKSNFQQ-----PYITNRTFMLEAKSLADNNNTDV 72
 Db 11 CLVLYLAPECGG----TDOG----DNFPOMLRLDRAFTSRVKTFFQTQD-----EVDN 56
 Qy 73 RLIGEKLFHGVMSRSRCYLMQKVNLNFTEBVLFPOSDFQPYMOSFVPPFLAR---LSNR 128
 Db 57 LLLKESLLEDFKGVLGQAIASEMIOFYLEEVM-FQAENQDPEAKDHVNSLGENLKTLR 115
 Qy 129 LSTCHIEGDDLHIGQNVQLRDTVKRLGSGGETAIGEIDL 169
 Db 116 LRRCHRFELCENKSKAQEIQKNAFNKLQFKGIYKAMSEFDI 156

Search completed: December 5, 2003, 18:21:31
 Job time : 29.64 secs

Result No.	Score	Query	Match	Length	DB ID	Description
1	903	100.0	179	21	AAB36292	Human GIL-19/AF289
2	903	100.0	179	21	AAT78081	Human T cell, induc
3	903	100.0	179	22	AAB04538	Human cytokine, ZC
4	903	100.0	179	22	AAB26664	Human IL-TIF Polyp
5	903	100.0	179	22	AAB77602	Human PRO10096. H
6	903	100.0	179	22	AB31210	Amino acid sequenc
7	903	100.0	179	22	AB48074	Human extracellular
8	903	100.0	179	23	AEE28608	Human IL-TIF prote
9	100.0	179	23		ABB79910	Human interleukin-

Pred. No. is the number of results predicted by a chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%	Query	Match	Length	DB ID	Description	
1	903	100.0	179	21	AAB36292	Human GIL-19/AF289
2	903	100.0	179	21	AAT78081	Human T cell, induc
3	903	100.0	179	22	AAB04538	Human cytokine, ZC
4	903	100.0	179	22	AAB26664	Human IL-TIF Polyp
5	903	100.0	179	22	AAB77602	Human PRO10096. H
6	903	100.0	179	22	AB31210	Amino acid sequenc
7	903	100.0	179	22	AB48074	Human extracellular
8	903	100.0	179	23	AEE28608	Human IL-TIF prote
9	100.0	179	23		ABB79910	Human interleukin-

Homo sapiens.

PN WO200065027-A2.

XX

PD 02-NOV-2000.

XX

PR 28-APR-2000; 2000WO-US11479.

XX

PR 28-APR-1998; 99US-0131473.

XX

PA (GEMY) GENETICS INST INC.

XX

PI Jacobs K, Fouser L, Spaulding V, Xuan D;

XX DR WPI: 2000-687325/67.

XX DR N-PSDB; AAC81773.

Human GIL-19 protein that shows a high degree of homology to IL

ALIGNMENTS

RESULT 1	AAB36292	standard; Protein; 179 AA.
ID		
XX		
AC	AAB36292;	
XX		
DT	23-FEB-2001 (First entry)	
XX		
DE	Human GIL-19/AF289 protein sequence.	
XX		
KW	Human; GIL-19/AF289; IL-10; interleukin-10; nutrition; cell proliferation; immune stimulation; tissue growth; inflammation; haemopoiesis regulation; tissue growth; inflammation; cancer.	
XX		
OS	Homo sapiens.	
XX		
PN	WO200065027-A2.	
XX		
PD	02-NOV-2000.	
XX		
PR	28-APR-2000; 2000WO-US11479.	
XX		
PR	28-APR-1998; 99US-0131473.	
XX		
PA	(GEMY) GENETICS INST INC.	
XX		
PI	Jacobs K, Fouser L, Spaulding V, Xuan D;	
XX		
DR	WPI: 2000-687325/67.	
XX		
DR	N-PSDB; AAC81773.	
XX		

PT (interleukin)-10, useful in upregulation of humoral immune responses, as an antiinflammatory agent and as a modulator of immune responses associated with injury -

PT Claim 9; Page 59-60; 60pp; English.

XX The present invention provides the protein and coding sequences for the novel human *IL-19/AAZ19* protein. The protein shows homology to interleukin-10 (IL-10) and is assumed to be a cytokine. It can be used in the regulation of cell proliferation and differentiation, hematopoiesis, immune stimulation or suppression, tissue growth and tumour inhibition. In addition, it also has uses in the treatment of inflammation and in nutrition.

XX Sequence 179 AA;

Query Match 100.0%; Score 903; DB 21; Length 179;
 Best Local Similarity 100.0%; Pred. No. 2.7e-86;
 Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAALQKVSSTSFNGTILATSCLLLLAVQGGAAPISHCRDLDSNFQQPYITNRTFMLA 60
 Db 1 MAALQKVSSTSFNGTILATSCLLLLAVQGGAAPISHCRDLDSNFQQPYITNRTFMLA 60

Qy 61 KEASLADNNNTDVLIGKLFGVMSRCYLMQKVINFTEBEVLFFQSDRQPYMQEVYP 120
 Db 61 KEASLADNNNTDVLIGKLFGVMSRCYLMQKVINFTEBEVLFFQSDRQPYMQEVYP 120

Qy 121 FLARLSNRSLSTCHIEGDDLHQIKNVQKLKDVTKLGESEGEIKAGELDLFMSLNACI 179
 Db 121 FLARLSNRSLSTCHIEGDDLHQIKNVQKLKDVTKLGESEGEIKAGELDLFMSLNACI 179

RESULT 2

AY92879 standard; Protein; 179 AA.
 XX
 AC AAY92879;
 XX
 DT 04-SEP-2000 (first entry)

XX Human T cell inducible factor.

XX TIF; T cell derived inducible factor; interleukin 9; STAT; IL-9;
 KW Anti-asthmatic; anti-allergic; cytostatic; inhibitor; antagonist;
 KW chromosome 12q15.
 XX Homo sapiens.

OS

XX

PD

04-MAY-2000.

XX

PN WO200024758-A1.

XX

PR 18-OCT-1999; 99WO-US24424.

XX

PR 26-OCT-1998; 98US-0178973.

XX

PR 16-JUL-1999; 99US-0354243.

XX

PA (LUDM-) LUDWIG INST CANCER RES.

XX

PA Dumoutier L, Louhed J, Renaud J;

XX

DR WPI; 2000-422495/36.

XX

PT New nucleic acid molecule encoding a T cell derived inducible factor

XX

PT for treating asthma, an allergy or lymphoma

XX

PS Example 26; Fig 1; 46pp; English.

XX

CC This is the human T cell derived inducible factor (TIF). The gene

CC was mapped to chromosome 12q15. The human TIF was identified based on

CC homology to a murine TIF, which was identified by subtraction cloning

CC from a murine lymphoma cell line BW5147 in the presence or absence of CC interleukin 9 (IL-9). BW5147 can be grown *in vitro*, without the need to CC add any cytokines to its culture medium. Many TIFs are CC mediated by activation of STAT transcription factors. The novel TIFs were CC expressed in the presence of IL-9, but not in its absence. TIFs induce CC STAT activation in cells. They can be used, e.g. in the stimulation of CC regeneration of targeted tissues. Their inhibitors or antagonists can be CC used to retard, prevent or inhibit differentiation of other tissues. The CC TIFs and their coding sequences are useful in the treatment of asthma, CC allergies and lymphoma (claimed). They are also useful for identifying CC compounds that inhibit or activate T cell induced factor activity in a CC cell (claimed).

XX Sequence 179 AA;

Query Match 100.0%; Score 903; DB 21; Length 179;
 Best Local Similarity 100.0%; Pred. No. 2.7e-86;
 Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAALQKVSSTSFNGTILATSCLLLLAVQGGAAPISHCRDLDSNFQQPYITNRTFMLA 60
 Db 1 MAALQKVSSTSFNGTILATSCLLLLAVQGGAAPISHCRDLDSNFQQPYITNRTFMLA 60

Qy 61 KEASLADNNNTDVLIGKLFGVMSRCYLMQKVINFTEBEVLFFQSDRQPYMQEVYP 120
 Db 61 KEASLADNNNTDVLIGKLFGVMSRCYLMQKVINFTEBEVLFFQSDRQPYMQEVYP 120

Qy 121 FLARLSNRSLSTCHIEGDDLHQIKNVQKLKDVTKLGESEGEIKAGELDLFMSLNACI 179
 Db 121 FLARLSNRSLSTCHIEGDDLHQIKNVQKLKDVTKLGESEGEIKAGELDLFMSLNACI 179

RESULT 3

AAE04538
 ID AAE04538 standard; Protein; 179 AA.

XX
 AC AAE04538;

DT 10-SEP-2001 (first entry)

XX Human cytokine, ZCYT018 protein #1.

XX Human; cytostatic; cytokine; ZCYT018 protein; genetic abnormality;

XX Cancer; inflammation; gene therapy.

XX Homo sapiens.

OS

XX

QY 121 FLARLSNRSLTICHIEGDDLHQIQRNVQKLKDTRVKLGESGETKAIGEGLDLIFMSLRNACI 179
 DB 121 FLARLSNRSLTICHIESDDLHQIQRNVQKLKDTRVKLGESGETKAIGEGLDLIFMSLRNACI 179

RESULT 7
 AAB48074 AAB48074 standard; protein; 179 AA.

XX XX

XX AAB48074;
 AC AC;
 XX XX

DT 19-MAR-2001 (first entry)

DE Human extracellular signaling molecule (EXCS) (ID 5571181CDI).

XX KW Extracellular signaling molecule; EXCS; anti-inflammatory; human;
 KW immunosuppressive; cytotoxic; neuroprotective; Gastrointestinal;
 KW virucide; antibacterial; anti-HIV; human immunodeficiency virus;
 KW antiinfection; cerebroprotective; nootropic; antiulcer; antitumor;
 KW anticonvulsant; tranquilizer; neuroleptic; vasoactive; gynecological;
 KW keratolytic; protozoacide; gene therapy.

OS Homo sapiens.

PN WO2000070049-A2.

XX XX

PD 23-NOV-2000.

PF 19-MAY-2000; 2000WO-US13975.

XX PR 19-MAY-1999; 99ITS-0114949.
 PR 15-JUL-1999; 99ITS-0114270.
 PR 30-JUL-1999; 99ITS-0146700.
 PR 04-OCT-1999; 99ITS-0137508.

XX PA (INCY-) INCYTE GENOMICS INC.

PI Tang YT, Yue H, Lal P, Burford N, Bandman O, Baughn MR;
 P1 Azimzai Y, Lu DAM, Patterson C;
 XX WPI; 2001-025021/03.
 DR N-PSDB; AAC84310.

PT New human extracellular signaling nucleic acids and polypeptides useful
 for diagnosing, treating and preventing infections and
 gastrointestinal, neurological, reproductive, and
 autoimmune/inflammatory disorders -

XX

PS Claim 1; Page 94; 114pp; English.

XX

CC The invention provides human extracellular signaling molecules (EXCS)
 CC and polynucleotides which identify and encode EXCS. EXCS can be
 CC expressed by standard recombinant methodology. The amino acid and nucleic
 CC acid sequences of EXCS are useful for diagnosing, treating and
 CC preventing infections and gastrointestinal (peptic ulcer, dysphagia,
 CC pancreatitis), neurological (e.g., epilepsy, ischemic cerebrovascular
 CC disease, stroke), reproductive (infertility, ovulatory defects, acquired
 CC immunodeficiency syndrome (AIDS), Addison's disease), and cell
 CC proliferative disorders including cancers (of the breast, adrenal gland,
 CC bone). They may also be used to treat fatal familial insomnia,
 CC nutritional and metabolic diseases of the nervous system, myopathies,
 CC mental disorders (anxiety, schizophrenia, mood), as well as infections
 CC caused by parasites (malaria, leishmania, trypanosoma), viral
 CC (adenovirus, coronavirus, flavivirus), bacterial (e.g., pneumococcus,
 CC staphylococcus, bacillus), and fungal (aspergillus, blastomycosis,
 CC dermatophytes) agents. The nucleic acids, polypeptides, antagonists,
 CC agonists, pharmaceutical compositions, and antibodies may also be used
 CC for treating or preventing disorders associated with increased or
 CC decreased expression or activity of EXCS. EXCS polynucleotides may also
 CC be used to detect and quantify gene expression in biopsied tissues in
 CC which expression of EXCS may be correlated with the disease, to determine

CC presence or excess expression of EXCS, to monitor regulation of EXCS
 CC levels during therapeutic intervention, to detect the presence of
 CC associated disorders, as targets in microarray, to generate hybridization
 CC probes, and to detect differences in gene sequences among normal, carrier
 CC or affected individuals. Antibodies may also be used in diagnosing, carrier
 CC disorders, in monitoring patients being treated with EXCS agonists,
 CC antagonists or inhibitors. Sequences AAB48057-B48082 represent the EXCS
 CC of the invention.

XX

SG Sequence 179 AA;

Query Match 100.0%; Score 903; DB 22; Length 179;
 Best Local Similarity 100.0%; Pred. No. 2.7e-86;
 Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAALQKSYSSFLNGTLLATSCLLLALLYQGAAAPISSHHCRLLDKSNFQPYITNRTFMLA 60
 DB 1 MAALQKSYSSFLNGTLLATSCLLLALLYQGAAAPISSHHCRLLDKSNFQPYITNRTFMLA 60

QY 61 KEASLADNNTDVRLIGELEFLHGYSMSERCYLMKQVLFNFTLEEVLFQSDRFQPYMOEVYP 120
 DB 61 KEASLADNNTDVRLIGELEFLHGYSMSERCYLMKQVLFNFTLEEVLFQSDRFQPYMOEVYP 120

QY 121 FLARLSNRSLTICHIEDDLDHQIQRNVQKLKDTRVKLGESGETKAIGEGLDLIFMSLRNACI 179
 DB 121 FLARLSNRSLTICHIEDDLDHQIQRNVQKLKDTRVKLGESGETKAIGEGLDLIFMSLRNACI 179

RESULT 8

AAE28608 ID AAE28608 standard; Protein, 179 AA.

XX AAE28608;
 AC AAE28608;
 XX DT 27-DEC-2002 (first entry)

XX Human IL-TIF protein #1.

DB Cytokine receptor; Zcytor16; IL-TIF; autoimmune disease; dermatological;
 KW inflammatory disease; inflammatory bowel disease; rheumatoid arthritis;
 KW asthma; systemic lupus erythematosus; myasthenia gravis; pancreatitis;
 KW diabetes; atherosclerosis; Glomerulonephritis; gene therapy; cytostatic;
 KW immuno suppressive; nephrotropic; allergy; placental health; abortion;
 KW cancer; human.

XX Homo sapiens.

XX WO200270655-A2.

XX 12-SEP-2002.

XX DF 04-MAR-2002; 2002WO-US06267.

XX PR 02-MAR-2001; 2001US-273035P.

XX PR 27-MAR-2001; 2001US-279232P.

XX PA (ZYMO) ZYMOGENETICS INC.

XX PI Presnell SR, Xu W, Kindsvogel W, Chen Z;

XX DR WPI; 2002-698750/75.

XX DR N-PSDB; AAD45964.

XX Disclosure: Page 197-198; 221pp; English.

XX New Zcytor16 polypeptide useful for treating autoimmune or inflammatory
 PT diseases, e.g. inflammatory bowel disease, rheumatoid arthritis,
 PT asthma, atherosclerosis, cancer or diabetes, or in assessing
 PT therapeutic aspects of IL-TIF -

XX

CC The invention relates to cytokine receptor designated as mouse Zcytor16
 CC which can bind and antagonize the IL-TIF. The Zcytor16 polypeptide is
 CC useful in modulating the immune system by binding Zcytor16 ligand, and

D 29-AUG-2002.
X 06-DEC-2001; 2001US-0006867.
F

polypeptide designated as A, B, C or D or E, F, G, H or I. The bioactive molecule is a toxin, a radiolabel or an antibody. The bioactive molecule causes death of the cell. A, B, C, D, E, F, G, H, or I, or antibodies against them are useful for modulating a biological activity of cell expressing a polypeptide designated as A, B, C or D or E, F, G, H, or I. The cell is killed. The Proteins are useful for identifying agonists or antagonists, for the preparation of a medicament useful in the treatment of a condition which is responsive to the proteins, as molecular weight markers for protein electrophoresis purposes, and as therapeutic agents for treating sports-related joint problems, articular cartilage defects, osteoarthritis or rheumatoid arthritis. Nucleic acids encoding the proteins are useful as hybridisation probes, in chromosome and gene mapping, in the generation of anti sense RNA and DNA, for the preparation of the proteins, to generate transgenic or knockout animals which are useful in the development and screening of therapeutic useful reagents. For chromosome identification, and in gene therapy. The antibody is useful as a therapeutic agent, in a diagnostic assay and for affinity purification of the protein from recombinant cell culture natural sources. The present sequence represents a novel secreted or transmembrane protein of the invention.

an extracellular domain of the proteins with their associated signal peptide or licking its associated signal peptide. Also included are the nucleic acids encoding the proteins, vectors, host cells, fusion proteins and antibodies which specifically bind to the proteins. The proteins are useful for detecting a polypeptide designated as A, B, C or D in a sample suspected of containing an A, B, C or D polypeptide, by contacting the sample with a polypeptide designated as E, F, G, H or I (or vice versa) and determining the formation of a A/E, B/F, C/G, C/H or D/I polypeptide conjugate in the sample, where the formation of the conjugate is indicative of the presence of an A, B, C or D polypeptide in the sample, where A is a PRO104272 polypeptide, B is a PRO20110 polypeptide, C is PRO10096 polypeptide, D is a PRO19760 polypeptide, E is a PRO5801 polypeptide, F is a PRO1 polypeptide, G is a PRO20040 polypeptide, H is a PRO20233 polypeptide and I is a PRO1890 polypeptide. The sample comprises a cell suspected of expressing the A, B, C or D polypeptide. The B, F, G, H or I polypeptide is labeled with a detectable label or is attached to a solid support. The proteins are useful for linking a bioactive molecule to a cell expressing a

CC disease (e.g. IDDM, multiple sclerosis (MS), systemic lupus erythematosus (SLE), myasthenia gravis, rheumatoid arthritis and IBD), CC sepsis, asthma, allergy and other atopic diseases, psoriasis and kidney dysfunction. Soluble zcytovII receptor or heterodimeric receptor polypeptides are useful in vivo or in diagnostic applications to detect CC IL-TF expressing cancers in vivo or in tissue samples and to prepare CC antibodies. ZcytovII serves as target for Mab therapy of cancer where CC an antagonising Mab inhibits cancer growth and targets immune-mediated CC killing. The present sequence represents the interleukin-22 (IL-22) CC inducible factor (IL-TIF) protein, the activity of this protein is CC inhibited by the zcytovII protein of the invention.

XX Sequence 179 AA;

Query Match 100.0%; Score 903; DB 23; Length 179;

Best Local Similarity 100.0%; Pred. No. 2.7e-86; Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MAALQKSVSSPLMGTATSCUILLALIVQGAAAAPISSHCFLDKSNFQOQYITNRTFMLA 60
 Db 1 MAALQKSVSSPLMGTATSCUILLALIVQGAAAAPISSHCFLDKSNFQOQYITNRTFMLA 60
 Db 61 KEASLADNNTIVRLIGEKLFFGVMSERCYLMQKVINFTEBVLPQSDRFQPMQEVV 120
 QY 61 KEASLADNNTIVRLIGEKLFFGVMSERCYLMQKVINFTEBVLPQSDRFQPMQEVV 120
 Db 61 KEASLADNNTIVRLIGEKLFFGVMSERCYLMQKVINFTEBVLPQSDRFQPMQEVV 120
 QY 121 FLARLSNRLSTKCHIEGDDLHIIQARNVQLKDTVKLGESGEKAIGELDILFMSLRNACI 179
 Db 121 FLARLSNRLSTKCHIEGDDLHIIQARNVQLKDTVKLGESGEKAIGELDILFMSLRNACI 179

RESULT

AAU78081 AAU78081 standard; Protein: 179 AA.

AC AAU78081;

DT 05-JUN-2002 (first entry)

XX Human interleukin 22 (IL-22) protein sequence.
 XX Interleukin 22; IL-22; cytostatic; antiinflammatory; IL-22 antagonist; KW immunotherapy; PAP1; pancreatic associated protein; receptor; receptor; KW IL-22R; IL-10beta; bioactive molecule linkage; cell death; pancreatitis; pancreatitis disorder; pancreatic carcinoma; acinar cell carcinoma; human; KW mixed cell population pancreatic carcinoma.

OS Homo sapiens.

XX Key

Peptide 1.3.3

FT Location/Qualifiers 1/label= Signal_peptide

FT Modified-site 14..20 /note= "Asn is N-myristoylated"

FT Protein 34..179 /label= Mature_human_interleukin_22_(IL-22)

FT Modified-site 54..58 /note= "Asn is N-glycosylated"

FT Modified-site 68..72 /note= "Asn is N-glycosylated"

FT Modified-site 82..88 /note= "Asn is N-myristoylated"

FT Modified-site 97..101 /note= "Asn is N-glycosylated"

XX WO200216611-A2.
 XX DE Human PRO10096 protein sequence SEQ ID NO:354.
 XX PD 28-FEB-2002.
 XX PP 30-MAY-2001; 2001WO-US17443.
 XX PR 24-AUG-2000; 2000WO-US23328.
 XX

PA (GETH) GENENTECH INC.

XX PI Aggarwal S, Foster JS, Goddard A, Gurney AL, Maruoka EM, Wood WI;

XX PI XIe M; DR WPI; 2002-28094/32.

XX DR N-PSDB; ABK11847.

XX Novel isolated interleukin 22 polypeptide useful for identifying IL-22 agonists and antagonists that are used for treating acute pancreatitis, chronic pancreatitis, pancreatic carcinoma -

XX SQ Claim 11; Fig 2; 94pp; English.

XX The present invention relates to a new polypeptide having at least 80% identity to a 101 amino acid interleukin (IL)-22 sequence. The invention is useful for detecting IL-22 (IL-22 receptor) or IL-10beta polypeptide in a sample which involves contacting simple with an IL-22 polypeptide and determining the formation of an IL-22/IL-22 polypeptide conjugate or polypeptide is labelled with a detectable label or is attached to a solid support. The polypeptide is also useful for linking a bioactive molecule, e.g. toxin, radiolabel or antibody that causes the death of the cell, to a cell expressing IL-22R polypeptide or IL-10beta polypeptide which involves contacting the cell with IL-22 polypeptide that is bound to the bioactive molecule and allowing binding of the IL-22 polypeptide with IL-22R or IL-10beta polypeptide thus linking the bioactive molecules to the cell. The molecules of the invention can also be used for modulating biological activity of cell expressing IL-22R or IL-10beta polypeptide, whereby the cell is killed and the antibody of the invention is useful for inhibiting IL-22 induced expression of PAP1 (pancreatitis associated protein) by pancreatic cells. The antibody is also useful for treating a pancreatic disorder such as acute or chronic pancreatitis, pancreatic carcinoma including acinar cell carcinoma or mixed cell population of the pancreas in a mammal. The present amino acid sequence represents the human interleukin 22 (IL-22) protein of the invention.

XX SO Sequence 179 AA;

XX Query Match 100.0%; Score 903; DB 23; Length 179;
 XX Best Local Similarity 100.0%; Pred. No. 2.7e-86;
 XX Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

XX QY 1 MAALQKSVSSPLMGTATSCUILLALIVQGAAAAPISSHCFLDKSNFQOQYITNRTFMLA 60
 XX Db 1 MAALQKSVSSPLMGTATSCUILLALIVQGAAAAPISSHCFLDKSNFQOQYITNRTFMLA 60
 XX QY 61 KEASLADNNTIVRLIGEKLFFGVMSERCYLMQKVINFTEBVLPQSDRFQPMQEVV 120
 XX Db 61 KEASLADNNTIVRLIGEKLFFGVMSERCYLMQKVINFTEBVLPQSDRFQPMQEVV 120
 XX QY 121 FLARLSNRLSTKCHIEGDDLHIIQARNVQLKDTVKLGESGEKAIGELDILFMSLRNACI 179
 XX Db 121 FLARLSNRLSTKCHIEGDDLHIIQARNVQLKDTVKLGESGEKAIGELDILFMSLRNACI 179
 XX QY 61 KEASLADNNTIVRLIGEKLFFGVMSERCYLMQKVINFTEBVLPQSDRFQPMQEVV 120
 XX Db 61 KEASLADNNTIVRLIGEKLFFGVMSERCYLMQKVINFTEBVLPQSDRFQPMQEVV 120
 XX QY 121 FLARLSNRLSTKCHIEGDDLHIIQARNVQLKDTVKLGESGEKAIGELDILFMSLRNACI 179
 XX Db 121 FLARLSNRLSTKCHIEGDDLHIIQARNVQLKDTVKLGESGEKAIGELDILFMSLRNACI 179
 XX RESULT 14
 XX ID ABB84993 standard; Protein: 179 AA.
 XX AC ABB84993;
 XX DT 16-MAY-2002 (first entry)

XX Human PRO10096 protein sequence SEQ ID NO:354.
 XX DE Human; angiogenesis; cardiotonic; cytostatic; antiangiogenic; hypotensive; KW vulnerability; antiatherosclerotic; PRO agonist; PRO antagonist; trauma; gene therapy; cardiovascular disorder; endothelial disorder; cancer; KW angiogenic disorder; cardiac hypertension; atherosclerosis; hypertension; KW age-related macular degeneration; arterial restenosis; angina; KW rheumatoid arthritis; myocardial infarction; thrombophlebitis;

KW lymphangitis; tumour angiogenesis; breast carcinoma; liver carcinoma;
 KW wound healing; chromosome mapping; gene mapping.

OS Homo sapiens.

XX WO200200530-A2.

XX 03-JAN-2002.

XX 20-JUN-2001; 2001WO-US19692.

XX 23-JUN-2000; 2000US-213637P.

PR 20-JUL-2000; 2000US-219556P.

PR 25-JUL-2000; 2000US-220524P.

PR 25-JUL-2000; 2000US-220664P.

PR 28-JUL-2000; 2000US-2206710.

PR 02-AUG-2000; 2000US-222995P.

PR 17-AUG-2000; 2000US-0641657.

PR 23-AUG-2000; 2000US-062522.

PR 07-SEP-2000; 2000US-0623328.

PR 18-SEP-2000; 2000US-062328P.

PR 24-OCT-2000; 2000US-0663350.

PR 08-NOV-2000; 2000US-0709238.

PR 10-NOV-2000; 2000US-0709352.

PR 01-DEC-2000; 2000US-0732678.

PR 20-DEC-2000; 2000US-0742759.

PR 22-FEB-2001; 2001US-0767609.

PR 28-FEB-2001; 2001US-0796498.

PR 28-FEB-2001; 2001WO-US06520.

PR 01-MAR-2001; 2001WO-US06666.

PR 09-MAR-2001; 2001US-0802706.

PR 14-MAR-2001; 2001US-0806839.

PR 05-APR-2001; 2001US-0816744.

PR 10-MAY-2001; 2001US-0828366.

PR 25-MAY-2001; 2001US-0854480.

PR 25-MAY-2001; 2001US-0866028.

PR 25-MAY-2001; 2001US-0866034.

PR 30-MAY-2001; 2001WO-US17092.

PR 30-MAY-2001; 2001US-0870574.

PR 01-JUN-2001; 2001WO-US17443.

PR 01-JUN-2001; 2001WO-US17800.

XX (GEETH) GENENTECH INC.

XX Baker KP, Ferrara N, Gerber H, Gerritsen ME, Goddard A, Godowski RJ, Gurney AL, Hillian KJ, Marsters SA, Pan J, Paoni NF, Stetefeld JF, Watanabe CK, Williams PM, Wood WI, Ye W,

XX WPI; 2002-090516/12.

XX DR; 2002-090516/12.

XX PS; ABL88248.

XX PT

PT One hundred and eighty seven nucleic acids encoding PRO polypeptides, useful in diagnosis and treatment of cardiovascular (e.g. myocardial infarction), endothelial or angiogenic disorders in a mammal -

XX XX

CC in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping; ABL88259 to ABL88667 represent primers and probes used in the exemplification of the present invention.

CC Sequence 179 AA;

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

XX SQ

CC Query Match 100.0%; Score 903; DB 23; Length 179;
 CC Best Local Similarity 100.0%; Pred. No. 2.7e-36;
 CC Matches 0; Mismatches 0; Indels 0; Gaps 0;

</div

Qy	1	MAALQKSVSSFLNGTLATSCLLILLALLVQGGAAPISSHCRLDKSNFQQPYITNRTEMLA	60
Db	1	MAALQKSVSSFLNGTLATSCLLILLALLVQGGAAPISSHCRLDKSNFQQPYITNRTEMLA	60
Qy	61	KEASLADNNTDYRLIGEKLFIGYMSERCYLMQKVINFITLEEVLFFQSDRQPYMOEVVP	120
Db	61	KEASLADNNTDYRLIGEKLFIGYMSERCYLMQKVINFITLEEVLFFQSDRQPYMOEVVP	120
Qy	121	FLARLSNRLSTCHIEGDDLH1QRNVQQLKDTVKKLGESEGEKAIGEGLLFLMSLRNACI	179
Db	121	FLARLSNRLSTCHIEGDDLH1QRNVQQLKDTVKKLGESEGEKAIGEGLLFLMSLRNACI	179

Search completed: December 5, 2003, 18:16:35
Job time : 61.9323 secs

121 FLARLSNRSLSTCHIEGDDLMIQRNVQKLKDTVKUIGESGBIAKGELDLEMIRNACI 179
Qy

PRIOR FILING DATE: 1999-05-19
PRIOR APPLICATION NUMBER: 60/144,270
PRIOR FILING DATE: 1999-07-15

RESULT 2
US-09-870-574-2
Sequence 2, Application US/09870574
Patent No. US20010102723A1
GENERAL INFORMATION:
APPLICANT: Gurney, Austin L.
APPLICANT: Aggarwal, Sudeepa
APPLICANT: Xie, Ming-Hong
APPLICANT: Maruoka, Ellen M.
APPLICANT: Foster, Jessica S.
APPLICANT: Goddard, Audrey
APPLICANT: Wood, William J.
TITLE OF INVENTION: INTERLEUKIN-22 POLYPEPTIDES, NUCLEIC ACIDS ENCODING
TITLE OF INVENTION: THE SAME AND METHODS FOR THE TREATMENT OF PANCREATIC DISORDERS
FILE REFERENCE: P2806-1 (US)
CURRENT APPLICATION NUMBER: US/09/870,574
CURRENT FILING DATE: 2001-05-30
PRIOR APPLICATION NUMBER: US 60/169,495
PRIOR FILING DATE: 1999-12-07
PRIOR APPLICATION NUMBER: PCT/US00/14042
PRIOR FILING DATE: 2000-05-22
PRIOR APPLICATION NUMBER: PCT/US00/23328
PRIOR FILING DATE: 2000-08-24
NUMBER OF SEQ ID NOS: 7

PRIOR APPLICATION NUMBER: 60/131,308
PRIOR FILING DATE: 1999-10-04
NUMBER OF SEQ ID NOS: 55

121 FLARLSNRSLSTCHIEGDDLHIQRNV

Query	Match	100.0%	Score	903	DB	10	Length	179	
Best Local	Similarity	100.0%	Pred.	No.	3..e-09				
Matches	179	Conservative	0	Mismatches	0	Indels	0	Gaps	0
Qy	1	MAALQKSVSSPIMGTLATSCILLALLIVQGAAAPISSHCRIDKSNFQQPYITNRTEMLA	60						
Db	1	MAALQKSVSSPLMGTIATSCILLALLIVQGAAATISSHCRIDKSNFQQPYITNRTEMLA	60						
Qy	61	KEASLADNNTDVRIGEKLPHGVNSERCYLMKQVNLNFTEEVLPQSDRFQPMQEVYP	120						
Db	61	KEASLADNNTDVRIGEKLPHGVNSERCYLMKQVNLNFTEEVLPQSDRFQPMQEVYP	120						
Qy	121	FLARLSNRLSTCHIEGDDLHQIQRNQKLKTVKLGESCEPIKAIGELDLMFLMSLRNACI	179						
Db	121	FLARLSNRLSTCHIEGDDLHQIQRNQKLKTVKLGESCEPIKAIGELDLMFLMSLRNACI	179						

US-10-063-735-154
; Sequence 154, Application US/10063735
; Publication No. US20030138882A1

Qy	61	XEASLADNNNDTVRLGERKLPHGVNSVSRCLMKQVNLNFTEEVIFPQSDRFQPMQEYPP	12.00
Db	61	KEASLADNNNDTVRLGERKLPHGVNSVSRCLMKQVNLNFTEEVIFPQSDRFQPMQEYPP	12.00
Qy	121	PLAISNLSTCHIEDDLLIQRNTQKLKDUTVKLGSEKIAICEDLFLMSRNACI	17.9
Db	121	FUABSNRVSSTCHIEDDLLIOPNTQKLKDUTVKLGSEKIAICEDLFLMSRNACI	17.9

APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.

RESULT 3
US -03-965-528-18

APPLICANT: WOOD, William L.
TITLE OF INVENTION: SECRETED AND TRA
TITLE OF INVENTION: ACIDS ENCODING
FILE REFERENCE: 523205167

PUBLICATION NO.: US2004167525A1
GENERAL INFORMATION:
APPLICANT: INCYTE GENOMICS, INC.
APPLICANT: TANG, Y. TOM
APPLICANT: YUE, Henry
APPLICANT: LAL, Preeti
APPLICANT: BURFORD, Neil
APPLICANT: BANDWAN, Olga
APPLICANT: BAUGHN, Mariah R.
APPLICANT: AZIMMAI, Valda
APPLICANT: LU, Duyang Aina M.
APPLICANT: PATERSON, Chandra
TITLE OF INVENTION: EXTRACELLULAR SIGNALING MOLECULES
FILE REFERENCE: PF-0701 USA
CURRENT AFFILIATION NUMBER: US/09/965,528
CURRENT FILING DATE: 2001-09-26

SEQ ID NO 154
LENGTH: 179
TYPE: PRT
ORGANISM: Homo sapiens

1 APPLICANT: LAL, Preeti
1 APPLICANT: BURFORD, Neil
1 APPLICANT: BANDIANI, Olga
1 APPLICANT: BAUGIN, Maria R.
1 APPLICANT: AZIMZAI, Yalda
1 APPLICANT: LU, Duyng Aina M.
1 APPLICANT: PATTERSON, Chandra
1 TITLE OF INVENTION: EXTRACELLULAR SIGNALING MOLECULES
1 FILE REFERENCE: PF-0701 USA
1 CURRENT APPLICATION NUMBER: US/09/965,528
1 CURRENT FILING DATE: 2001-09-26
1 PRIOR APPLICATION NUMBER: 60/134,949

Query Match 100.0% ; Score
Best Local Similarity 100.0% ; Predict

1 APPLICANT: AZIMAI, Yalda
1 ;
1 APPLICANT: LU, Duyng Aina M.
1 ;
1 APPLICANT: PATTERSON, Chandra
1 ;
1 TITLE OF INVENTION: EXTRACELLULAR SIGNALING MOLECULES
1 ;
1 FILE REFERENCE: PP-0701 USA
1 ;
1 CURRENT APPLICATION NUMBER: US/09/965,528
1 ;
1 CURRENT FILING DATE: 2001-09-26
1 ;

CURRENT APPLICATION NUMBER: US/09/965, 528
CURRENT FILING DATE: 20/11/09-26
PRIOR APPLICATION NUMBER: 60/134, 949

Qy 61 KEASLADNNTDVRLIGEKLFLFGVSM
Db 61 KEASLADNNTDVRLIGEKLFLFGVSM

Qy 121 FLARLSNLSTCHIEGDDLHIIQRNVQKLDIVKLGESGRKIAIGEGLDLMFLMSRNACI 179
 Db 121 FLARLSNLSTCHIEGDDLHIIQRNVQKLDIVKLGESGRKIAIGEGLDLMFLMSRNACI 179

RESULT 5
 US-10-216-163-244
 Sequence 244. Application US/10216163
 Publication No. US20030149239A1
 GENERAL INFORMATION:
 APPLICANT: Baker, Kevin P.
 APPLICANT: Desnoyers, Luc
 APPLICANT: Gerritsen, Mary
 APPLICANT: Goddard, Audrey
 APPLICANT: Grimaldi, J. Christopher
 APPLICANT: Gurney, Austin L.
 APPLICANT: Smith, Victoria
 APPLICANT: Stephan, Jean-Philippe F.
 APPLICANT: Watansabe, Colin L.
 APPLICANT: Wood, William I.

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 FILE REFERENCE: P3530PLC3

CURRENT APPLICATION NUMBER: US 10/216,163
 CURRENT FILING DATE: 2002-08-09
 PRIOR APPLICATION NUMBER: 10/119,480
 PRIOR FILING DATE: 2002-04-09
 PRIOR APPLICATION NUMBER: 60/059,113
 PRIOR FILING DATE: 1997-09-17
 PRIOR APPLICATION NUMBER: 60/062,287
 PRIOR FILING DATE: 1997-10-17
 PRIOR APPLICATION NUMBER: 60/063,549
 PRIOR FILING DATE: 1997-10-28
 PRIOR APPLICATION NUMBER: 60/064,103
 PRIOR FILING DATE: 1997-10-31
 PRIOR APPLICATION NUMBER: 60/069,873
 PRIOR FILING DATE: 1997-12-17
 PRIOR APPLICATION NUMBER: 60/078,910
 PRIOR FILING DATE: 1998-03-20
 PRIOR APPLICATION NUMBER: 60/079,294
 PRIOR FILING DATE: 1998-03-25
 PRIOR APPLICATION NUMBER: 60/079,556
 PRIOR FILING DATE: 1998-03-26
 PRIOR APPLICATION NUMBER: 60/079,728
 PRIOR FILING DATE: 1998-03-27
 Remaining Prior Application data removed - See File Wrapper or PALM.
 SEQ ID NO: 246
 LENGTH: 179
 ORGANISM: Homo Sapien

Query Match 100.0%; Score 903; DB 12; Length 179;
 Best Local Similarity 100.0%; Pred. No. 3.1e-90; Indels 0; Gaps 0;
 Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAALQKSVSSFLMGTATSCLLALLVQGAAAPISSHCRDLSKNSFOQPYITNRTFMLA 60
 Db 1 MAALQKSVSSFLMGTATSCLLALLVQGAAAPISSHCRDLSKNSFOQPYITNRTFMLA 60

Qy 61 KEASLADNTDYLIGEKLFGVSMBSRCYLMKQVINFITLEEVLFQSDRQPTMQEYVP 120
 Db 61 KEASLADNTDYLIGEKLFGVSMBSRCYLMKQVINFITLEEVLFQSDRQPTMQEYVP 120

Qy 121 FLARLSNLSTCHIEGDDLHIIQRNVQKLDIVKLGESGRKIAIGEGLDLMFLMSRNACI 179
 Db 121 FLARLSNLSTCHIEGDDLHIIQRNVQKLDIVKLGESGRKIAIGEGLDLMFLMSRNACI 179

RESULT 6
 US-10-216-163-244
 Sequence 244. Application US/10216163
 Publication No. US20030149239A1
 GENERAL INFORMATION:
 APPLICANT: Topouris, Stavros
 FILE REFERENCE: 00-56
 CURRENT APPLICATION NUMBER: US/09/925,055D
 CURRENT FILING DATE: 2001-08-08
 PRIOR APPLICATION NUMBER: US 60/223,827
 PRIOR FILING DATE: 2000-08-08
 PRIOR APPLICATION NUMBER: US 60/250,876
 PRIOR FILING DATE: 2000-12-01
 NUMBER OF SEQ ID NOS: 35
 SOFTWARE: FASTSEQ for Windows Version 3.0
 SEQ ID NO: 8
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo sapiens
 US-09-925-055D-8

Query Match 100.0%; Score 903; DB 12; Length 179;

Best Local Similarity 100.0%; Pred. No. 3.1e-90;
 Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAALQKSVSSFLMGTATSCLLALLVQGAAAPISSHCRDLSKNSFOQPYITNRTFMLA 60
 Db 1 MAALQKSVSSFLMGTATSCLLALLVQGAAAPISSHCRDLSKNSFOQPYITNRTFMLA 60

Qy 61 KEASLADNTDYLIGEKLFGVSMBSRCYLMKQVINFITLEEVLFQSDRQPTMQEYVP 120
 Db 61 KEASLADNTDYLIGEKLFGVSMBSRCYLMKQVINFITLEEVLFQSDRQPTMQEYVP 120

Qy 121 FLARLSNLSTCHIEGDDLHIIQRNVQKLDIVKLGESGRKIAIGEGLDLMFLMSRNACI 179
 Db 121 FLARLSNLSTCHIEGDDLHIIQRNVQKLDIVKLGESGRKIAIGEGLDLMFLMSRNACI 179

Query Match 100.0%; Score 903; DB 12; Length 179;

Best Local Similarity 100.0%; Pred. No. 3.1e-90; Mismatches 0; Indels 0; Gaps 0; Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAALQKSVSSFLMGTATSCLLLALLVQGAAAPISSHCRDLDSNFQPYTNRTMFLA 60
Db 1 MAALQKSVSSFLMGTATSCLLLALLVQGAAAPISSHCRDLDSNFQPYTNRTMFLA 60

Qy 61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKVQVNFTLEEVLFQSDRFQPYMCEVVP 120
Db 61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKVQVNFTLEEVLFQSDRFQPYMCEVVP 120

Qy 121 FLARLSNRSLSTCHIEGDDLHIIQRNVQKLKDTVKLGSEGEIAKGELDLFMSLRNACI 1.79
Db 121 FLARLSNRSLSTCHIEGDDLHIIQRNVQKLKDTVKLGSEGEIAKGELDLFMSLRNACI 1.79

RESULT 8
US-09-7446-375-2
Sequence 2, Application US/097446375
; Publication No. US20030170823A1
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
; APPLICANT: Kindvogel, Wayne
; TITLE OF INVENTION: NOVEL CYTOKINE ZCYT018
; FILE REFERENCE: 99-106
; CURRENT APPLICATION NUMBER: US/09/746,375
; CURRENT FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 60/172,105
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: US 60/****,***
; NUMBER OF SEQ ID NOS: 44
; SOFTWARE: FastSSQ for Windows Version 3.0
; SEQ ID NO: 2
; LENGTH: 179
; TYPE: PT
; ORGANISM: Homo sapiens
US-09-7446-375-2

Query Match 100.0%; Score 903; DB 12; Length 179;
Best Local Similarity 100.0%; Pred. No. 3.1e-90; Mismatches 0; Indels 0; Gaps 0; Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MAALQKSVSSFLMGTATSCLLLALLVQGAAAPISSHCRDLDSNFQPYTNRTMFLA 60
Db 1 MAALQKSVSSFLMGTATSCLLLALLVQGAAAPISSHCRDLDSNFQPYTNRTMFLA 60

Qy 61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKVQVNFTLEEVLFQSDRFQPYMCEVVP 120
Db 61 KEASLADNNTDVRLIGEKLFHGVSMSERCYLMKVQVNFTLEEVLFQSDRFQPYMCEVVP 120

Qy 121 FLARLSNRSLSTCHIEGDDLHIIQRNVQKLKDTVKLGSEGEIAKGELDLFMSLRNACI 1.79
Db 121 FLARLSNRSLSTCHIEGDDLHIIQRNVQKLKDTVKLGSEGEIAKGELDLFMSLRNACI 1.79

RESULT 10
US-10-066-198-126
Sequence 126, Application US/10066198
; Publication No. US20030170721A1
; GENERAL INFORMATION:
; APPLICANT: Avi J. Ashkenazi
; APPLICANT: Kevin P. Baker
; APPLICANT: David A. Botein
; APPLICANT: Luc Desnoyers
; APPLICANT: Dan L. Eaton
; APPLICANT: Napoleon Ferrara
; APPLICANT: Sherman Fong
; APPLICANT: Wei-Qiang Gao
; APPLICANT: Hanspeter Gerber
; APPLICANT: Mary E. Gerritsen
; APPLICANT: Paul J. Godowski
; APPLICANT: Austin L. Gurney
; APPLICANT: Ivar J. Klijavins
; APPLICANT: Jennie P. Mather
; APPLICANT: Mary A. Napiers
; APPLICANT: James Pan
; APPLICANT: Nicholas P. Paonni
; APPLICANT: Margaret Ann Roy
; APPLICANT: Timothy A. Stewart
; APPLICANT: Daniel Tumas
; APPLICANT: Colin K. Watanabe
; APPLICANT: P. Mickey Williams
; APPLICANT: William I. Wood
; APPLICANT: Zemin Zhang

; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; ACIDS ENCODING THE SAME
; FILE REFERENCE: P3130R1C6

; CURRENT APPLICATION NUMBER: US/10/066,198
; PRIOR APPLICATION NUMBER: 10/002,796
; PRIOR FILING DATE: 2002-02-01
; PRIOR APPLICATION NUMBER: 10/002,796
; PRIOR FILING DATE: 1997-08-26
; PRIOR APPLICATION NUMBER: 50/059115
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/059263
; PRIOR FILING DATE: 1997-09-18
; PRIOR APPLICATION NUMBER: 60/059588
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/062285

RESULT 9
US-10-063-126-154
Sequence 154, Application US/10063526
; Publication No. US20030171550A1
; GENERAL INFORMATION:
; APPLICANT: Eaton, Dan L.
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Gerritsen, Mary E.
; APPLICANT: Goddard, Audrey
; APPLICANT: Grimaldi, Paul J.
; APPLICANT: Grimaldi, Christopher J.
; APPLICANT: Gurney, Austin L.
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Wood, William I.
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; ACIDS ENCODING THE SAME
; FILE REFERENCE: P3230R1C1

PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/062816
PRIOR FILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063082
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/063329
PRIOR FILING DATE: 1997-10-27
PRIOR APPLICATION NUMBER: 60/063733
PRIOR FILING DATE: 1997-10-29
PRIOR APPLICATION NUMBER: 60/065364
PRIOR FILING DATE: 1997-11-21
PRIOR APPLICATION NUMBER: 60/065840
PRIOR FILING DATE: 1997-11-25
PRIOR APPLICATION NUMBER: 60/065944
PRIOR FILING DATE: 1997-12-16
PRIOR APPLICATION NUMBER: 60/074086
PRIOR FILING DATE: 1998-02-09
PRIOR APPLICATION NUMBER: 60/074092
PRIOR FILING DATE: 1998-02-09
PRIOR APPLICATION NUMBER: 60/074092
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/081049
PRIOR FILING DATE: 1998-04-08
PRIOR APPLICATION NUMBER: 60/095998
PRIOR FILING DATE: 1998-08-10
PRIOR APPLICATION NUMBER: 60/097000
PRIOR FILING DATE: 1998-08-18
PRIOR APPLICATION NUMBER: 60/099601
PRIOR FILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: 60/099803
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099811
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099812
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/100858
PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: 60/101922
PRIOR FILING DATE: 1998-09-24
PRIOR APPLICATION NUMBER: 60/100322
PRIOR FILING DATE: 1998-10-28
PRIOR APPLICATION NUMBER: 60/10304
PRIOR FILING DATE: 1998-11-20
PRIOR APPLICATION NUMBER: 60/125778
PRIOR FILING DATE: 1999-03-23
PRIOR APPLICATION NUMBER: 60/133695
PRIOR FILING DATE: 1999-06-15
PRIOR APPLICATION NUMBER: 60/145070
PRIOR FILING DATE: 1999-07-20
PRIOR APPLICATION NUMBER: 60/145698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: 60/14996
PRIOR FILING DATE: 1999-08-17
PRIOR APPLICATION NUMBER: 60/163495
PRIOR FILING DATE: 1999-12-07
PRIOR APPLICATION NUMBER: 08/913874
PRIOR FILING DATE: 1999-08-16
PRIOR APPLICATION NUMBER: 08/933821
PRIOR FILING DATE: 1999-09-19
PRIOR APPLICATION NUMBER: 08/960507
PRIOR FILING DATE: 1999-10-29
PRIOR APPLICATION NUMBER: 09/114844
PRIOR FILING DATE: 1999-07-14
PRIOR APPLICATION NUMBER: 09/134601
PRIOR FILING DATE: 1999-08-19
PRIOR APPLICATION NUMBER: 09/136804
PRIOR FILING DATE: 1999-08-19
PRIOR APPLICATION NUMBER: 09/136828
PRIOR FILING DATE: 1999-08-19
PRIOR APPLICATION NUMBER: 09/153342
PRIOR FILING DATE: 1999-09-21
PRIOR APPLICATION NUMBER: 09/180997
PRIOR FILING DATE: 1999-09-10

1 PRIOR FILING DATE: 1999-03-08
 1 PRIOR APPLICATION NUMBER: PCT/US99/122252
 1 PRIOR FILING DATE: 1999-06-02
 1 PRIOR APPLICATION NUMBER: PCT/US99/20111
 1 PRIOR FILING DATE: 1999-09-01
 1 PRIOR APPLICATION NUMBER: PCT/US99/20594
 1 PRIOR FILING DATE: 1999-09-08
 1 PRIOR APPLICATION NUMBER: PCT/US99/21090
 1 PRIOR FILING DATE: 1999-09-15
 1 PRIOR APPLICATION NUMBER: PCT/US99/21547

Query Match Score 903; DB 12; Length 179;
 Best Local Similarity 100.0%; Pred. No. 3.1e-90;
 Matches 179; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1
 CURRENT APPLICATION NUMBER: US/10/063,510
 CURRENT FILING DATE: 2002-05-01
 PRIOR Application removed - See File Wrapper or Palm
 NUMBER OF SEQ ID NOS: 170
 SEQ ID NO 154
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-063-510-154

Query Match Score 903; DB 12; Length 179;
 Best Local Similarity 100.0%; Pred. No. 3.1e-90;
 Matches 179; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1
 CURRENT APPLICATION NUMBER: US/10/063,510
 CURRENT FILING DATE: 2002-05-01
 PRIOR Application removed - See File Wrapper or Palm
 NUMBER OF SEQ ID NOS: 170
 SEQ ID NO 154
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-063-510-154

RESULT 11
 US-10-063-586-154
 Sequence 154, Application US/10063586
 Publication No. US20030176684A1
 GENERAL INFORMATION:
 1 APPLICANT: Eaton, Dan L.
 1 APPLICANT: Filvaroff, Ellen
 1 APPLICANT: Gerritsen, Mary E.
 1 APPLICANT: Goddard, Audrey
 1 APPLICANT: Godowski, Paul J.
 1 APPLICANT: Grimaldi, Christopher J.
 1 APPLICANT: Gurney, Austin L.
 1 APPLICANT: Watanabe, Colin K.
 1 APPLICANT: Wood, William I.
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1
 CURRENT FILING DATE: 2002-05-03
 PRIOR Application removed - See File Wrapper or Palm
 NUMBER OF SEQ ID NOS: 170
 SEQ ID NO 154
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-063-586-154

Query Match Score 903; DB 12; Length 179;
 Best Local Similarity 100.0%; Pred. No. 3.1e-90;
 Matches 179; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1
 CURRENT APPLICATION NUMBER: US/10/063,514
 CURRENT FILING DATE: 2002-05-01
 PRIOR Application removed - See File Wrapper or Palm
 NUMBER OF SEQ ID NOS: 170
 SEQ ID NO 154
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-063-514-154

Query Match Score 903; DB 12; Length 179;
 Best Local Similarity 100.0%; Pred. No. 3.1e-90;
 Matches 179; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1
 CURRENT APPLICATION NUMBER: US/10/063,514
 CURRENT FILING DATE: 2002-05-01
 PRIOR Application removed - See File Wrapper or Palm
 NUMBER OF SEQ ID NOS: 170
 SEQ ID NO 154
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-063-514-154

Query Match Score 903; DB 12; Length 179;
 Best Local Similarity 100.0%; Pred. No. 3.1e-90;
 Matches 179; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1
 CURRENT APPLICATION NUMBER: US/10/063,514
 CURRENT FILING DATE: 2002-05-01
 PRIOR Application removed - See File Wrapper or Palm
 NUMBER OF SEQ ID NOS: 170
 SEQ ID NO 154
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-063-514-154

Query Match Score 903; DB 12; Length 179;
 Best Local Similarity 100.0%; Pred. No. 3.1e-90;
 Matches 179; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1
 CURRENT APPLICATION NUMBER: US/10/063,514
 CURRENT FILING DATE: 2002-05-01
 PRIOR Application removed - See File Wrapper or Palm
 NUMBER OF SEQ ID NOS: 170
 SEQ ID NO 154
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-063-514-154

Query Match Score 903; DB 12; Length 179;
 Best Local Similarity 100.0%; Pred. No. 3.1e-90;
 Matches 179; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1
 CURRENT APPLICATION NUMBER: US/10/063,514
 CURRENT FILING DATE: 2002-05-01
 PRIOR Application removed - See File Wrapper or Palm
 NUMBER OF SEQ ID NOS: 170
 SEQ ID NO 154
 LENGTH: 179
 TYPE: PRT
 ORGANISM: Homo Sapien
 US-10-063-514-154

RESULT 12

Y	1	MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISSHCRLDKSNFQQPYITNRTMLA	60
Y	1	MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISSHCRLDKSNFQQPYITNRTMLA	60
Y	61	KPASLADNNTDVRLIGEKLPHGYSMSPRCLMVKQVINFTEELVLFPSDRFQPYMQEVVP	120
Y	61	KPASLADNNTDVRLIGEKLPHGYSMSPRCLMVKQVINFTEELVLFPSDRFQPYMQEVVP	120
Y	121	FLARLSNRLSTCHIEGDLHICRNQVKLKDITVKLGSSEIKAGELDLMSLRNACI	179
Y	121	FLARLSNRLSTCHIEGDLHICRNQVKLKDITVKLGSSEIKAGELDLMSLRNACI	179
RESULT 14			
S-10-063-516-154			
Sequence 154, Application US/10063516			
Publication No. US20030181708A1			
GENERAL INFORMATION:			
APPLICANT:	Baton, Dan L.		
APPLICANT:	Filvaroff, Ellen		
APPLICANT:	Gerritsen, Mary E.		
APPLICANT:	Goddard, Audrey		
APPLICANT:	Godowski, Paul J.		
APPLICANT:	Grimaldi, Christopher J.		
APPLICANT:	Gurney, Austin L.		
APPLICANT:	Watanabe, Colin K.		
APPLICANT:	Wood, William I.		
TITLE OF INVENTION: ACIDS ENCODING THE SAME			
FILE REFERENCE: P2230RLC1			
CURRENT APPLICATION NUMBER: US/10/063_516			
CURRENT FILING DATE: 2002-05-01			
Prior Application removed - See File Wrapper or Palm			
NUMBER OF SEQ ID NOS: 170			
SEQ ID NO 154			
LENGTH: 179			
TYPE: PRT			
ORGANISM: Homo Sapien			
S-10-063-516-154			
Query Match 100.0%; Score 903; DB 12;			
Best Local Similarity 100.0%; Pred. No. 3.1e-50;			
Matches 179; Conservative 0; Mismatches 0; Indels 0; Gaps 0			
1 MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISSHCRLDKSNFQQPYITNRTMLA			
1 MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISSHCRLDKSNFQQPYITNRTMLA			
61 KEPASLADNNTDVRLIGEKLPHGYSMSPRCLMVKQVINFTEELVLFPSDRFQPYMQEVVP			
61 KEPASLADNNTDVRLIGEKLPHGYSMSPRCLMVKQVINFTEELVLFPSDRFQPYMQEVVP			
121 FLARLSNRLSTCHIEGDLHICRNQVKLKDITVKLGSSEIKAGELDLMSLRNACI			
121 FLARLSNRLSTCHIEGDLHICRNQVKLKDITVKLGSSEIKAGELDLMSLRNACI			

ES-15
S-10-063-523-154
Sequence 154, Application US/10063523
Publication No. US20030181636A1
GENERAL INFORMATION:
APPLICANT: Baton, Dan L.
APPLICANT: Filveroff, Ellen
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC

1 MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISHCRDKSNFQQPYITINRTFMLA 60
 1 MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISHCRDKSNFQQPYITINRTFMLA 60

61 KEASLADNNNTDVLIGKLFQGYSMERSCYLMKVQVNFTEBEVLFFQSDRQPYQEVVP 120
 61 KEASLADNNNTDVLIGKLFQGYSMERSCYLMKVQVNFTEBEVLFFQSDRQPYQEVVP 120

121 FLARLSRNLSTCHIEGDDLHQIQRNQVKKLDVKLGESGEIKAGIDLLEMISLNAI 179
 121 FLARLSRNLSTCHIEGDDLHQIQRNQVKKLDVKLGESGEIKAGIDLLEMISLNAI 179

IS-10-063-516-154
 Sequence 154, Application US/10063516
 Publication No. US20030181708A1
 GENERAL INFORMATION:
 APPLICANT: Eaton,Dan L.
 APPLICANT: Filvaroff,Ellen
 APPLICANT: Gerrissen,Mary E.
 APPLICANT: Goddard,Audrey
 APPLICANT: Godowski,Paul J.
 APPLICANT: Grimaldi,Christopher J.
 APPLICANT: Gurney,Austin L.
 APPLICANT: Watanabe,Colin K.
 APPLICANT: Wood,William I.
 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
 ACIDS ENCODING THE SAME
 FILE REFERENCE: P3230R1C1

1 MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISHCRDKSNFQQPYITINRTFMLA 60
 1 MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISHCRDKSNFQQPYITINRTFMLA 60

61 KEASLADNNNTDVLIGKLFQGYSMERSCYLMKVQVNFTEBEVLFFQSDRQPYQEVVP 120
 61 KEASLADNNNTDVLIGKLFQGYSMERSCYLMKVQVNFTEBEVLFFQSDRQPYQEVVP 120

121 FLARLSRNLSTCHIEGDDLHQIQRNQVKKLDVKLGESGEIKAGIDLLEMISLNAI 179
 121 FLARLSRNLSTCHIEGDDLHQIQRNQVKKLDVKLGESGEIKAGIDLLEMISLNAI 179

Qy 1 MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISHCRDKSNFQQPYITINRTFMLA 60
 Db 1 MAALQKSVSSPMLGTLATSCLLALLIVQGAAAPISHCRDKSNFQQPYITINRTFMLA 60

Qy 61 KEASLADNNNTDVLIGKLFQGYSMERSCYLMKVQVNFTEBEVLFFQSDRQPYQEVVP 120
 Db 61 KEASLADNNNTDVLIGKLFQGYSMERSCYLMKVQVNFTEBEVLFFQSDRQPYQEVVP 120

Qy 121 FLARLSRNLSTCHIEGDDLHQIQRNQVKKLDVKLGESGEIKAGIDLLEMISLNAI 179
 Db 121 FLARLSRNLSTCHIEGDDLHQIQRNQVKKLDVKLGESGEIKAGIDLLEMISLNAI 179

Search completed: December 5, 2003, 18:32:11
 Job time: 346.883 secs

Search completed: December 5, 2003, 18:32:11
Job time : 346.883 secs